

0590

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/054,399

Source: 01PE

Date Processed by STIC: 9-16-5

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Does Not Comply Corrected Diskette Needed



OIPE

RAW SEQUENCE LISTING DATE: 09/16/2002 PATENT APPLICATION: US/10/054,399 TIME: 10:54:29

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Output Set: N:\CRF4\09162002\J054399.raw

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             Nern, Peter MA
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      7 <130> FILE REFERENCE: DYOU13.1A2CP1
      9 <140> CURRENT APPLICATION NUMBER: 10/054399
C--> 10 <141> CURRENT FILING DATE: 2002-08-30
    12 <150> PRIOR APPLICATION NUMBER: 09/168,474
    13 <151> PRIOR FILING DATE: 1998-10-08
    15 <150> PRIOR APPLICATION NUMBER: 08/951,141
    16 <151> PRIOR FILING DATE: 1997-10-15
    18 <150> PRIOR APPLICATION NUMBER: 09/529,106
    19 <151> PRIOR FILING DATE: 2000-04-07
    21 <150> PRIOR APPLICATION NUMBER: US 09/732,180
    22 <151> PRIOR FILING DATE: 2000-12-07
    24 <150> PRIOR APPLICATION NUMBER: US 60/169,699
    25 <151> PRIOR FILING DATE: 1999-12-07
    27 <150> PRIOR APPLICATION NUMBER: PCT/GB98/03033
    28 <151> PRIOR FILING DATE: 1998-10-08
    30 <150> PRIOR APPLICATION NUMBER: 9812793.9
    31 <151> PRIOR FILING DATE: 1998-06-12
    33 <150> PRIOR APPLICATION NUMBER: 9721357.3
    34 <151> PRIOR FILING DATE: 1997-10-08
    36 <150> PRIOR APPLICATION NUMBER: 9721358.1
    37 <151> PRIOR FILING DATE: 1997-10-08
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    46 <213> ORGANISM: Saccharomyces cerevisiae
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    53 gacgatttga aagtotgtaa aaaatocatt tatgacttta tattgggotg caagaaacac 120
    54 tttgcattta acgatgagga gettttcact atatecgaeg tttttgccaa etcgaegtee 180
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DATE: 09/16/2002

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PATENT APPLICATION: US/10/054,399
                                                           TIME: 10:54:29
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                 Output Set: N:\CRF4\09162002\J054399.raw
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                                                                the Sequence Listing. Please check subsequent
65 <400> SEQUENCE: 2
                                                                sequences for similar errors.
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67
69 Val Ile Ala Ser Asp Asp Leu Lys Val Cys Lys Lys Ser Ile Tyr Asp
70
                 20
                                      25
72 Phe Ile Leu Gly Cys Lys Lys His Phe Ala Phe Asn Asp Glu Glu Leu
                                  40
75 Phe Thr Ile Ser Asp Val Phe Ala Asn Ser Thr Ser Gln Leu Val Lys
                                                                      need to explain the Source of the genetic Imaterial
76.
                              55
                                                  60
78 Val Leu Glu Val Val Glu Thr Leu Met Asn Ser Ser
79 65
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92 gacgatttga aagtetgtaa aaaateeatt tatgaettta tattgggetg caagaaacae 120
93 tttgcattta acgatgagga gcttttcact atatccgacg tttttgccaa ctcgacgtcc 180
94 cagctggtca aagtgctaga agtagtagaa acgctaatga attccagc
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109
                  20
111 Phe Ile Leu Gly Cys Lys Lys His Phe Ala Phe Asn Asp Glu Glu Leu
             35
112
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114 Phe Thr Ile Ser Asp Val Phe Ala Asn Ser Thr Ser Gln Leu Val Lys
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117 Val Leu Glu Val Val Glu Thr Leu Met Asn Ser Ser
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132 tttgcattta acgatgagga gettiteact atateegacg tttttgccaa etegacgtee 180

RAW SEQUENCE LISTING

DATE: 09/16/2002

TIME: 10:54:29

Input Set : A:\pto_ms.txt Output Set: N:\CRF4\09162002\J054399.raw 133 cagctggtca aagtgctaga agtagtagaa acgctaatga attccagc 228 135 <210> SEQ ID NO: 6 136 <211> LENGTH: 76 137 <212> TYPE: PRT 138 <213> ORGANISM: Artificial Sequence 140 <220> FEATURE: 141 <223> OTHER INFORMATION: Description of Artificial Sequence: amino acid 143 <400> SEQUENCE: 6 144 Pro Leu Cys Ile Leu Phe Asn Ser Val Lys Pro Gln Phe Lys Leu Pro 145 147 Val Ile Ala Ser Gly Asp Leu Lys Val Cys Lys Lys Ser Ile Tyr Asp 148 20 25 150 Phe Ile Leu Gly Cys Lys Lys His Phe Ala Phe Asn Asp Glu Glu Leu 151 35 45 153 Phe Thr Ile Ser Asp Val Phe Ala Asn Ser Thr Ser Gln Leu Val Lys 55 154 50 156 Val Leu Glu Val Val Glu Thr Leu Met Asn Ser Ser 157 65 160 <210> SEQ ID NO; 7 161 <211> LENGTH: 228 162 <212> TYPE: DNA 163 <213> ORGANISM: Artificial Sequence 165 <220> FEATURE: 166 <223> OTHER INFORMATION: Description of Artificial Sequence: nucleic acid 168 <400> SEQUENCE: 7 169 cccctctgta tacttttcaa ctctgtgaag ccgcaattta aattaccggt aatagcacct 60 170 gacgatttga aagtotgtaa aaaatooatt tatgaottta tattgggotg caagaaacac 120 171 ttttgcattta acgatgagga gettttcaet atatecgaeg tttttgccaa etegaegtee 180 172 cagotggtoa aagtgctaga agtagtagaa acgctaatga attccago 174 <210> SEQ ID NO: 8 175 <211> LENGTH: 76 176 <212> TYPE: PRT 177 <213> ORGANISM: Artificial Sequence 179 <220> FEATURE: 180 <223> OTHER INFORMATION: Description of Artificial Sequence: amino acid 182 <400> SEQUENCE: 8 183 Pro Leu Cys Ile Leu Phe Asn Ser Val Lys Pro Gln Phe Lys Leu Pro 184 1 186 Val Ile Ala Pro Asp Asp Leu Lys Val Cys Lys Lys Ser Ile Tyr Asp 187 189 Phe Ile Leu Gly Cys Lys Lys His Phe Ala Phe Asn Asp Glu Glu Leu 190 35 4.0 192 Phe Thr Ile Ser Asp Val Phe Ala Asn Ser Thr Ser Gln Leu Val Lys 55 195 Val Leu Glu Val Val Glu Thr Leu Met Asn Ser Ser 196 65 70

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/054,399

199 <210> SEQ ID NO: 9 200 <211> LENGTH: 392 201 <212> TYPE: PRT

see payed

RAW SEQUENCE LISTING PATENT APPLICATION: US/10/054,399

DATE: 09/16/2002 TIME: 10:54:29

Input Set : A:\pto_ms.txt

Output Set: N:\CRF4\09162002\J054399.raw

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211 Leu Glu Ile Leu Asp Lys Tyr Arg Gln Gln Leu Leu Asp Ser Asn Leu
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214 Ile Thr Ser Glu Glu Leu Tyr Met Leu Phe Pro Asn Leu Gly Asp Ala
            35
                                40
217 Ile Asp Phe Gln Arg Arg Phe Leu Ile Ser Leu Glu Ile Asn Ala Leu
       50
                            55
218
220 Val Glu Pro Ser Lys Gln Arg Ile Gly Ala Leu Phe Met His Ser Lys
221 65
                        ..70
                                            75
223 His Phe Phe Lys Leu Tyr Glu Pro Trp Ser Ile Gly Gln Asn Ala Ala
                    85
                                        90
226 Ile Glu Phe Leu Ser Ser Thr Leu His Lys Met Arg Val Asp Glu Ser
               100
                                   105
                                                       110
229 Gln Arg Phe Ile Ile Asn Asn Lys Leu Glu Leu Gln Ser Phe Leu Tyr
                                                   125
           115
                               120
230
232 Lys Pro Val Gln Arg Leu Cys Arg Tyr Pro Leu Leu Val Lys Glu Leu
                           1.35
                                               140
233
       130
235 Leu Ala Glu Ser Ser Asp Asp Asn Asn Thr Lys Glu Leu Glu Ala Ala
                                           155
                       150
238 Leu Asp Ile Ser Lys Asn Ile Ala Arg Ser Ile Asn Glu Asn Gln Arg
                   165
                                       170
                                                           175
241 Arg Thr Glu Asn His Gln Val Val Lys Lys Leu Tyr Gly Arg Val Val
               180
                                   185
                                                       190
242
244 Asn Trp Lys Gly Tyr Arg Ile Ser Lys Phe Gly Glu Leu Leu Tyr Phe
245
                               200
                                                   205
           195
247 Asp Lys Val Phe Ile Ser Thr Thr Asn Ser Ser Ser Glu Pro Glu Arg
                                               220
                           215
       210
250 Glu Phe Glu Val Tyr Leu Phe Glu Lys Ile Ile Ile Leu Phe Ser Glu
251 225
                       230
                                           235
253 Val Val Thr Lys Lys Ser Ala Ser Ser Leu Ile Leu Lys Lys Ser
                   245
                                       250
256 Ser Thr Ser Ala Ser Ile Ser Ala Ser Asn Ile Thr Asp Asn Asn Gly
                                                       270
                                   265
257
                260
259 Ser Pro His His Ser Tyr His Lys Arg His Ser Asn Ser Ser Ser Ser
                                                   285
           275
                               280
260
262 Asn Asn Ile His Leu Ser Ser Ser Ala Ala Ala Ile Ile His Ser
                           295
263 290
265 Ser Thr Asn Ser Ser Asp Asn Asn Ser Asn Asn Ser Ser Ser Ser Ser
                                           315
                       310
268 Leu Phe Lys Leu Ser Ala Asn Glu Pro Lys Leu Asp Leu Arg Gly Arg
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                   325
269
271 Ile Met Ile Met Asn Leu Asn Gln Ile Ile Pro Gln Asn Asn Arg Ser
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272
274 Leu Asn Ile Thr Trp Glu Ser Ile Lys Glu Gln Gly Asn Phe Leu Leu
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DATE: 09/16/2002

TIME: 10:54:29

Input Set : A:\pto_ms.txt Output Set: N:\CRF4\09162002\J054399.raw 360 355 277 Lys Phe Lys Asn Glu Glu Thr Arg Asp Asn Trp Ser Ser Cys Leu Gln 375 380 278 370: 280 Gln Leu Ile His Asp Leu Lys Asn 281 385 390 284 <210> SEQ ID NO: 10 285 <211> LENGTH: 1269 286 <212> TYPE: DNA 287 <213> ORGANISM: Saccharomyces cerevisiae 289 <220> FEATURE: 290 <223> OTHER INFORMATION: 292 <400> SEQUENCE: 10 293 atggcacate agatggacte gataacgtat tetaataatg teacceaaca gtatatacaa 60 294 ccacaaagto tacaggatat ctotgcagtg gaggaagaaa ttoaaaataa aatagaggoo 120 295 gccagacaag agagtaaaca gcttcatgct caaataaata aagcaaaaca caagatacaa 180 296 gatgcaaget tattecagat ggecaacaaa gttaettegt tgaccaaaaa taagatcaac 240 297 ttaaagccaa atatcgtgtt gaaaggccat aataataaaa tctcagattt tcggtggagt 300 298 cgagattcaa aacgtatttt gagtgcaagt caagatggct ttatgcttat atgggacagt 360 299 getteaggtt taaaacagaa egetatteea ttagattete aatgggttet tteetgeget 420 300 atttcgccat cgagtacttt ggtagcaagc gcaggattaa acaataactg taccatttat 480 301 agagtttega aagaaaacag agtagegeaa aaegttgegt caatttteaa aggacataet 540 302 tyctatattt ctgacattga atttacagat aacgcacata tattgacagc aagtggggat 600 303 atgacatgtg cottgtggga tatacogaaa gcaaagaggg tgagagaata ttotgaccat 660 304 thaggtgatg tittggcatt agctattect gaagageeaa acttagaaaa ticttegaac 720 305 acattogota gotgtggato agacgggtat acttacatat gggatagoag atotocgtoc 780 306 gctgtacaaa gcttttacgt taacgatagt gatattaatg cacttcgttt tttcaaagac 840 307 gggatgtega ttgttgcagg aagtgacaat ggtgegataa atatgtatga tttaaggteg 900 308 gactgttcta ttgctacttt ttctcttttt cgaggttatg aagaacgtac ccctacccct 960 309 acttatatgg cagctaacat ggagtacaat accgcgcaat cgccacaaac tttaaaatca 1020 310 acaageteaa getatetaga caaceaagge gttgtttett tagattttag tgcatetgga 1080 311 agattgatgt actcatgcta tacagacatt ggttgtgttg tgtgggatgt attaaaagga 1140 312 gagattgttg gaaaattaga aggtcatggt ggcagagtca ctggtgtgcg ctcgagtcca 1200 313 gatgggttag ctgtatgtac aggttcatgg gactcaacca tgaaaatatg gtctccaggt 1260 314 tatcaatag 316 <210> SEQ ID NO: 11 317 <211> LENGTH: 422 318 <212> TYPE: PRT 319 <213> ORGANISM: Saccharomyces cerevisiae 321 <220> FEATURE: 322 <223> OTHER INFORMATION: 324 <400> SEQUENCE: 11 325 Met Ala His Gln Met Asp Ser Ile Thr Tyr Ser Asn Asn Val Thr Gln 328 Gln Tyr Ile Gln Pro Gln Ser Leu Gln Asp Ile Ser Ala Val Glu Glu 329 20 25 331 Glu Ile Gln Asn Lys Ile Glu Ala Ala Arg Gln Glu Ser Lys Gln Leu

40

334 His Ala Gln Ile Asn Lys Ala Lys His Lys Ile Gln Asp Ala Ser Leu

55

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/054,399

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35

332

335

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/054,399

DATE: 09/16/2002 TIME: 10:54:30

Input Set : A:\pto_ms.txt

Output Set: N:\CRF4\09162002\J054399.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:32; N Pos. 12 Seq#:33; N Pos. 13 VERIFICATION SUMMARY
PATENT APPLICATION: US/10/054,399

DATE: 09/16/2002 TIME: 10:54:30

Input Set : A:\pto_ms.txt

Output Set: N:\CRF4\09162002\J054399.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:1799 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0 L:1816 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0 L:1843 M:283 W: Missing Blank Line separator, <400> field identifier

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/054,399

DATE: 09/10/2002 TIME: 08:57:09

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Output Set: N:\CRF3\09102002\J054399.raw

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L:1841 M:282 W: Numeric Field Identifier Missing, <213> is required.

6:1841 M:200 E: Mandatory Header Field missing, <400> is required.
6:39 M:203 E: No. of Seq. differs, <160> Number Of Sequences: Input (37) Counted (36)

Does Not Comply Corrected Diskette Needed



OTPE

RAW SEQUENCE LISTING

DATE: 09/10/2002

PATENT APPLICATION: US/10/054,399

TIME: 08:57:08

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Output Set: N:\CRF3\09102002\J054399.raw

- 2 <110> APPLICANT: Arkowitz, Robert A
- Nern, Peter MA
- 5 <120> TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND PROTEIN SEQUENCES
- 7 <130> FILE REFERENCE: DYOU13.1A2CP1
- 9 <140> CURRENT APPLICATION NUMBER: 10/054399
- C--> 10 <141> CURRENT FILING DATE: 2002-08-30
 - 12 <150> PRIOR APPLICATION NUMBER: 09/168,474
 - 13 <151> PRIOR FILING DATE: 1998-10-08
 - 15 <150> PRIOR APPLICATION NUMBER: 08/951,141
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 - 18 <150> PRIOR APPLICATION NUMBER: 09/529,106
 - 19 <151> PRIOR FILING DATE: 2000-04-07
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 - 24 <150> PRIOR APPLICATION NUMBER: US 60/169,699
 - 25 <151> PRIOR FILING DATE: 1999-12-07
 - 27 <150> PRIOR APPLICATION NUMBER: PCT/GB98/03033
 - 28 <151> PRIOR FILING DATE: 1998-10-08
 - 30 <150> PRIOR APPLICATION NUMBER: 9812793.9
 - 31 <151> PRIOR FILING DATE: 1998-06-12
 - 33 <150> PRIOR APPLICATION NUMBER: 9721357.3
 - 34 <151> PRIOR FILING DATE: 1997-10-08
 - 36 <150> PRIOR APPLICATION NUMBER: 9721358.1
 - 37 <151> PRIOR FILING DATE: 1997-10-08
- -> 39 <160> NUMBER OF SEQ ID NOS: 37
 - 41 <170> SOFTWARE: PatentIn Ver. 2.0

need to insert Line breaks between numeric identifiers.

ERRORED SEQUENCES

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- W--> 1841 <212> TYPE:
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1842 1 5

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